RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	09/518 2970
Source:	IFW/6,
Date Processed by STIC:	7/7/06

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 07/07/2006 PATENT APPLICATION: US/09/518,297D TIME: 17:17:03

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\1518297D.raw

```
4 <110> APPLICANT: Lim, Moon Young
      5 Edwards, Cynthia A.
            Fry, Kirk E.
      7
            Bruice, Thomas W.
             Starr, Douglas B.
      9
             Laurance, Megan E.
         Kwok, Yan
     10
     13 <120> TITLE OF INVENTION: DNA Binding Compound-Mediated Molecular
     14 Switch System
     16 <130> FILE REFERENCE: 4600-0130.30
     18 <140> CURRENT APPLICATION NUMBER: US 09/518,297D
     19 <141> CURRENT FILING DATE: 2000-03-03
     21 <150> PRIOR APPLICATION NUMBER: US 60/122,513
     22 <151> PRIOR FILING DATE: 1999-03-03
     24 <150> PRIOR APPLICATION NUMBER: US 60/154,605
     25 <151> PRIOR FILING DATE: 1999-09-17
     27 <160> NUMBER OF SEQ ID NOS: 77
     29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     31 <210> SEO ID NO: 1
     32 <211> LENGTH: 11
     33 <212> TYPE: DNA
     34 <213> ORGANISM: Artificial Sequence
     36 <220> FEATURE:
     37 <223> OTHER INFORMATION: DNA response element
     39 <400> SEQUENCE: 1
     40 cgttcgcact t
                                                                               11
     42 <210> SEO ID NO: 2
     43 <211> LENGTH: 17
     44 <212> TYPE: DNA
     45 <213> ORGANISM: Artificial Sequence
     47 <220> FEATURE:
     48 <223> OTHER INFORMATION: DNA response element
     50 <400> SEQUENCE: 2
     51 cggagtactg tcctccg
                                                                               17
     53 <210> SEQ ID NO: 3
     54 <211> LENGTH: 12
     55 <212> TYPE: DNA
     56 <213> ORGANISM: Artificial Sequence
     58 <220> FEATURE:
     59 <223> OTHER INFORMATION: DNA response element
W--> 61 <221> NAME/KEY: misc feature
    62 <222> LOCATION: (1)...(12)
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63 <223 > OTHER INFORMATION: n = A, T, C or G

RAW SEQUENCE LISTING DATE: 07/07/2006 PATENT APPLICATION: US/09/518,297D TIME: 17:17:03

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\I518297D.raw

```
W--> 65 <400> 3
W--> 66 taattanggg ng
                                                                               12
     68 <210> SEQ ID NO: 4
     69 <211> LENGTH: 551
     70 <212> TYPE: PRT
     71 <213> ORGANISM: Homo sapiens
     73 <220> FEATURE:
     74 <221> NAME/KEY: VARIANT
     75 <222> LOCATION: (0)...(0)
     76 <223> OTHER INFORMATION: transcriptional regulatory protein
     78 <400> SEQUENCE: 4
    79 Met Asp Glu Leu Phe Pro Leu Ile Phe Pro Ala Glu Pro Ala Gln Ala
                        5
     81 Ser Gly Pro Tyr Val Glu Ile Ile Glu Gln Pro Lys Gln Arg Gly Met
                    20
     83 Arg Phe Arg Tyr Lys Cys Glu Gly Arg Ser Ala Gly Ser Ile Pro Gly
     85 Glu Arg Ser Thr Asp Thr Thr Lys Thr His Pro Thr Ile Lys Ile Asn
     87 Gly Tyr Thr Gly Pro Gly Thr Val Arg Ile Ser Leu Val Thr Lys Asp
                            70
    89 Pro Pro His Arg Pro His Pro His Glu Leu Val Gly Lys Asp Cys Arg
                                            90
    91 Asp Gly Phe Tyr Glu Ala Glu Leu Cys Pro Asp Arg Cys Ile His Ser
                    100
                                        105
    93 Phe Gln Asn Leu Gly Ile Gln Cys Val Lys Lys Arg Asp Leu Glu Gln
               115
                                    120
                                                        125
    95 Ala Ile Ser Gln Arg Ile Gln Thr Asn Asn Pro Phe Gln Val Pro
                                135
    97 Ile Glu Glu Gln Arg Gly Asp Tyr Asp Leu Asn Ala Val Arg Leu Cys
                           150
                                                155
    99 Phe Gln Val Thr Val Arg Asp Pro Ser Gly Arg Pro Leu Arg Leu Pro
                        165
    101 Pro Val Leu Pro His Pro Ile Phe Asp Asn Arg Ala Pro Asn Thr Ala
                    180
                                         185
    103 Glu Leu Lys Ile Cys Arg Val Asn Arg Asn Ser Gly Ser Cys Leu Gly
                                    200
    105 Gly Asp Glu Ile Phe Leu Leu Cys Asp Lys Val Gln Lys Glu Asp Ile
                                215
                                                     220
    107 Glu Val Tyr Phe Thr Gly Pro Gly Trp Glu Ala Arg Gly Ser Phe Ser
                            230
                                                 235
    109 Gln Ala Asp Val His Arg Gln Val Ala Ile Val Phe Arg Thr Pro Pro
                        245
                                             250
    111 Tyr Ala Asp Pro Ser Leu Gln Ala Pro Val Arg Val Ser Met Gln Leu
                                         265
    113 Arg Arg Pro Ser Asp Arg Glu Leu Ser Glu Pro Met Glu Phe Gln Tyr
                                     280
    115 Leu Pro Asp Thr Asp Asp Arg His Arg Ile Glu Glu Lys Arg Lys Arg
```

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RAW SEQUENCE LISTING DATE: 07/07/2006 PATENT APPLICATION: US/09/518,297D TIME: 17:17:03

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\I518297D.raw

		Tyr	Glu	Thr	Phe		Ser	Ile	Met	Lys	_	Ser	Pro	Phe	Ser	_		
	305	m1	•	n	_	310		_	_	_	315			_	_	320		
	Pro	Thr	Asp	Pro		Pro	Pro	Pro	Arg		Ile	Ala	Val	Pro		Arg		
120	_	_			325				_	330	_				335			
	Ser	Ser	Ala	Ser	Val	Pro	Lys	Pro	Ala	Pro	Gln	Pro	Tyr	Pro	Phe	Thr		
122				340					345					350				
123	Ser	Ser	Leu	Ser	Thr	Ile	Asn	Tyr	Asp	Glu	Phe	Pro	Thr	Met	Val	Phe		
124			355					360					365					
125	Pro	Ser	Gly	Gln	Ile	Ser	Gln	Ala	Ser	Ala	Leu	Ala	Pro	Ala	Pro	Pro		
126		370					375					380						
127	Gln	Val	Leu	Pro	Gln	Ala	Pro	Ala	Pro	Ala	Pro	Ala	Pro	Ala	Met	Val		
	385					390					395					400		
		Ala	Leu	Ala	Gln		Pro	Ala	Pro	Val		Val	Len	Δla	Pro			
130					405					410					415	017		
	Pro	Pro	Gln	Δla		Δla	Pro	Pro	Δla		Lve	Dro	Thr	Gln		Glv		
132	110	110	0111	420	vai	nia	110	110	425	110	цуз	110	1111	430	лта	Gry		
	Clu	G1 v	Thr		C02	C1,,	Ala	T 011		C1 n	Tou	C1 n	Dho		7	C1		
	GIU	Gry	435	ьец	ser	GIU	AIA		neu	GIII	ьeu	GIII		Asp	Asp	GIU		
134	7	T		77-	T	.	~ 1	440	0	m1			445		51	m1		
	Asp		GIY	Ala	ьeu	ьeu	Gly	ASI	Ser	Thr	Asp		Ата	vaı	Pne	Inr		
136		450		_		_	455	_		_,		460	_	_	_			
		Leu	Ата	ser	Val		Asn	Ser	GIu	Phe		Gin	Leu	Leu	Asn			
	465			_	_	470				_	475					480		
	Gly	Ile	Pro	Val		Pro	His	Thr	Thr	Glu	Pro	Met	Leu	Met		Tyr		
140					485					490					495			
141	Pro	Glu	Ala		Thr	Arg	Leu	Val	Thr	Gly	Ala	Gln	Arg	Pro	Pro	Asp		
142				500					505					510				
143	Pro	Ala	Pro	Ala	Pro	Leu	Gly	Ala	Pro	Gly	Leu	Pro	Asn	Gly	Leu	Leu		
144			515					520					525					
145	Ser	Gly	Asp	Glu	Asp	Phe	Ser	Ser	Ile	Ala	Asp	Met	Asp	Phe	Ser	Ala		
146		530					535					540						
147	Leu	Leu	Ser	Gln	Ile	Ser	Ser											
148	545					550												
150	<210)> SI	EQ II	NO:	5													
		L> LE																
152	<212	2> T)	PE:	DNA														
153	<213	3> OF	(GAN	SM:	Arti	fici	ial S	Seque	ence									
								•										
	5 <220> FEATURE: 5 <223> OTHER INFORMATION: DNA response element																	
)> SE				-			<u>r</u>									
			-			a												19
	9 tccctatcag tgatagaga 1 <210> SEQ ID NO: 6																	
	2 <211> LENGTH: 22																	
	2 <211> DENGIH: 22 3 <212> TYPE: DNA																	
	4 <213> ORGANISM: Artificial Sequence																	
	6 <220> FEATURE:																	
	56 <220> FEATURE: 57 <223> OTHER INFORMATION: response element																	
						TOM:	. res	POUR	oc e.	LemeI	ıL							
)> SE					_											22
		acac				.a aç	d											22
1/2	<210)> SE	יל דד	NO:	,													

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DATE: 07/07/2006

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/518,297D TIME: 17:17:03

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\1518297D.raw

```
173 <211> LENGTH: 13
     174 <212> TYPE: DNA
    175 <213> ORGANISM: Artificial Sequence
    177 <220> FEATURE:
     178 <223> OTHER INFORMATION: response element
W--> 180 <221> NAME/KEY: misc feature
     181 <222> LOCATION: (3)...(3)
    182 <223> OTHER INFORMATION: n = G or T
W--> 184 <221> misc feature
    185 <222> LOCATION: (7)...(7)
    186 <223> OTHER INFORMATION: n = A, T, C or G
W--> 188 <221> misc feature
    189 <222> LOCATION: (12)...(12)
    190 <223> OTHER INFORMATION: n = A or C
W--> 192 <400> 7
W--> 193 rgntcantga cny
    195 <210> SEQ ID NO: 8
    197 <400> SEQUENCE: 8
W--> 198 000
    200 <210> SEQ ID NO: 9
    202 <400> SEQUENCE: 9
W--> 203 000
    205 <210> SEQ ID NO: 10
    206 <211> LENGTH: 97
    207 <212> TYPE: PRT
    208 <213> ORGANISM: Artificial Sequence
    210 <220> FEATURE:
    211 <223> OTHER INFORMATION: repressor sequence
    213 <400> SEQUENCE: 10
    214 Met Asp Ala Lys Ser Leu Thr Ala Trp Ser Arg Thr Leu Val Thr Phe
                                             10
    216 Lys Asp Val Phe Val Asp Phe Thr Arg Glu Glu Trp Lys Leu Leu Asp
                     20
    218 Thr Ala Gln Gln Ile Val Tyr Arg Asn Val Met Leu Glu Asn Tyr Lys
                                     40
    220 Asn Leu Val Ser Leu Gly Tyr Gln Leu Thr Lys Pro Asp Val Ile Leu
                                 55
    222 Arg Leu Glu Lys Gly Glu Glu Pro Trp Leu Val Glu Arg Glu Ile His
                            70
                                                 75
    224 Gln Glu Thr His Pro Asp Ser Glu Thr Ala Phe Glu Ile Lys Ser Ser
    225
                                             90
    226 Val
    229 <210> SEQ ID NO: 11
    230 <211> LENGTH: 36
    231 <212> TYPE: PRT
    232 <213> ORGANISM: Artificial Sequence
    234 <220> FEATURE:
    235 <223> OTHER INFORMATION: repressor sequence
    237 <400> SEQUENCE: 11
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RAW SEQUENCE LISTING DATE: 07/07/2006 PATENT APPLICATION: US/09/518,297D TIME: 17:17:03

Input Set: A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\1518297D.raw

238 Met Ala Ala Ala Val Arg Met Asn Ile Gln Met Leu Leu Glu Ala Ala 239 1 3.0 240 Asp Tyr Leu Glu Arg Arg Glu Arg Glu Ala Glu His Gly Tyr Ala Ser 241 242 Met Leu Pro Tyr 243 35 245 <210> SEQ ID NO: 12 246 <211> LENGTH: 116 247 <212> TYPE: DNA 248 <213> ORGANISM: Escherichia coli 250 <220> FEATURE: 251 <221> NAME/KEY: misc feature 252 <222> LOCATION: (0)...(0) 253 <223> OTHER INFORMATION: partial promoter sequence 255 <400> SEQUENCE: 12 256 cgcggtcaga aaattatttt aaatttcctc ttgtcaggcc ggaataactc cctataatgc 60 257 gccaccactg acacggaaca acggcaaaca cgccgccqqq tcaqcqqqqt tctcct 116 259 <210> SEQ ID NO: 13 260 <211> LENGTH: 22 261 <212> TYPE: DNA 262 <213> ORGANISM: Escherichia coli 264 <220> FEATURE: 265 <221> NAME/KEY: misc_feature 266 <222> LOCATION: (0)...(0) 267 <223> OTHER INFORMATION: partial promoter sequence 269 <400> SEQUENCE: 13 270 agaaaattat tttaaatttc ct 22 272 <210> SEQ ID NO: 14 273 <211> LENGTH: 22 274 <212> TYPE: DNA 275 <213> ORGANISM: Artificial Sequence 277 <220> FEATURE: 278 <223> OTHER INFORMATION: modified promoter sequence 280 <400> SEQUENCE: 14 281 gactgcagtg gtacctagga gg 22 283 <210> SEQ ID NO: 15 284 <211> LENGTH: 22 285 <212> TYPE: DNA 286 <213> ORGANISM: Artificial Sequence 288 <220> FEATURE: 289 <223> OTHER INFORMATION: modified promoter sequence 291 <400> SEQUENCE: 15 292 agaaaattat tttaaatttc ct 22 294 <210> SEQ ID NO: 16 295 <211> LENGTH: 22 296 <212> TYPE: DNA 297 <213> ORGANISM: Artificial Sequence 299 <220> FEATURE: 300 <223> OTHER INFORMATION: modified promoter sequence

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/07/2006
PATENT APPLICATION: US/09/518,297D TIME: 17:17:04

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\1518297D.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 7,11 Seq#:7; N Pos. 3,7,12 VERIFICATION SUMMARYDATE: 07/07/2006PATENT APPLICATION: US/09/518,297DTIME: 17:17:04

Input Set : A:\54600-8130.US00-SEQLIST.TXT
Output Set: N:\CRF4\07072006\I518297D.raw

L:61 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:65 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:180 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:184 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:188 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:192 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:198 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (8) SEQUENCE:
L:203 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (63) SEQUENCE:
L:1211 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (64) SEQUENCE: